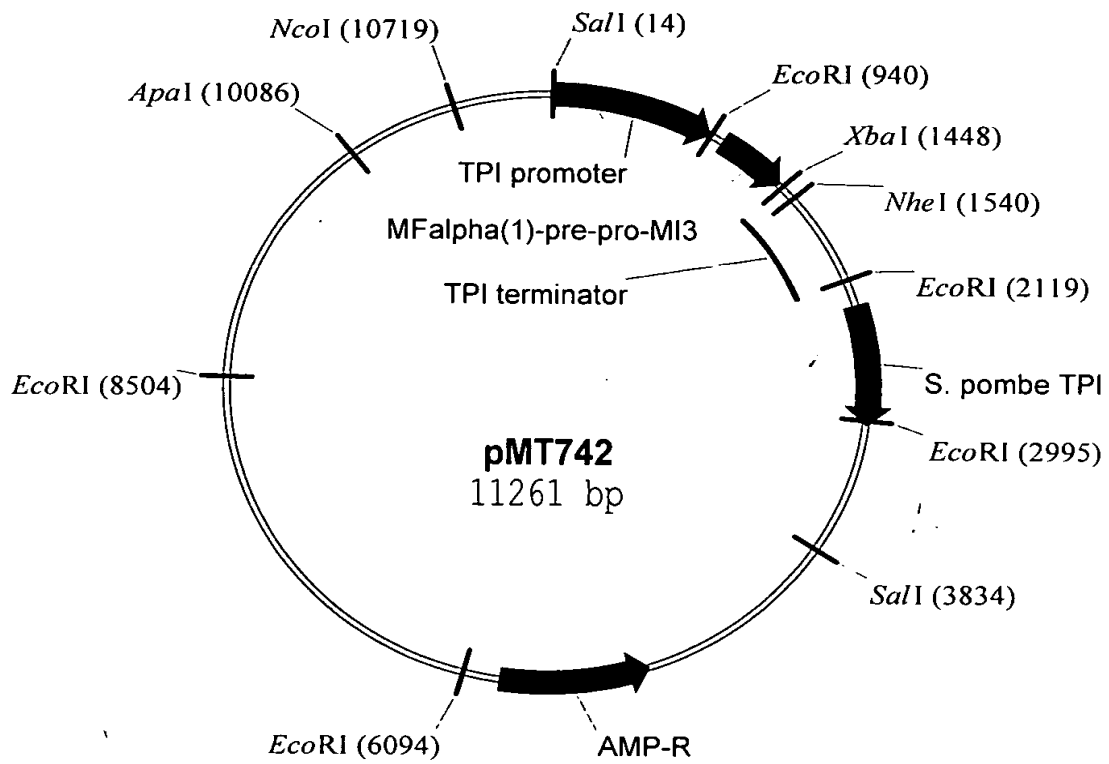


Fig. 1



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Fig. 2

5

G GA

1 TATAAGAGAG CTCATCTTAT TGTTGTCAGC CCAATGATTC CCTTGTCAAA
51 TTGAATTTTC GGATTTACTT GTTCAGGTAC CCGCGTTAAG GGGCTGCCGC
101 GCCTGTCACT CTAAGAAAAA AGGAGCCATC AAAAACCATT CAGCATTAAC
10 151 TAAAAACGCG GGTAGAGATT ACTACATATT CCAACAAGAC CTTCGCAGGA
201 AAGTATACCT AACTAATTA AAGAAATCTC CGAAGTTCGC ATTTTCATTGA
251 ACGGCTCAAT TAATCTTTGT AAATATGAGC GTTTTTACGT TCACATTGCC
301 TTTTTTTTTT TGTATTTACC TTGCATTTTT GTGCTAAAAG GCGTCACGTT
351 TTTTTCCGCC GCAGCCGCC GGAAATGAAA AGTATGACCC CCGCTAGACC
15 401 AAAAATACTT TTGTGTTATT GGAGGATCGC AATCCCTTG GAGCTTTTCC
451 GATACTATCG ACTTATCCGA CCTCTTGTTG TTTGAAAATG TCAATTGATA
501 TCCATCCATT ATATAAATGC TCAAACTTG CAGCAACTAT TCTTTACCCT
551 TCCCCTGTTA TGGATTGCTA GTCTTAAGGG GGAAATTGTC TGTTTACTAA
601 AATACAAACC AGGTTTGTTT TGGCTTTTAT TTGCATTTAA GTAATTACAA
20 651 TTACAACCAT TAAAAAGAAA ATAAGGCAAA ACATATAGCA ATATAATACT
701 ATTTACGAAG ATGTCAGCGA TA

T TC